



<110> Glaxo Group Limited  
Gauthier, Jean-Michel

<120> Method of screening

<130> PF3402

<140> PCT/EP99/00654

<141> 1999-02-04

<150> GB 9802475.5

<151> 1998-02-06

<160> 23

<170> PatentIn Ver. 2.1

<210> 1

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
construct

<400> 1

agccagacaa gccagacaag ccagacaagc cagacaagcc agacaagcca gacaagccag 60  
acaagccaga caagccagac a 81

<210> 2

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
construct

<400> 2

agacagacaa gacagacaag acagacaaga cagacaagac agacaagaca gacaagacag 60  
acaagacaga caagacagac a 81

<210> 3

<211> 81

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic  
construct

<400> 3

agctacataa gctacataag ctacataagc tacataagct acataagcta cataagctac 60  
ataagctaca taagctacat a 81

<210> 4

<211> 467

<212> PRT

<213> Homo sapiens

<400> 4

Met Ser Ser Ile Leu Pro Phe Thr Pro Pro Val Val Lys Arg Leu Leu  
1 5 10 15

Gly Trp Lys Lys Ser Ala Gly Gly Ser Gly Gly Ala Gly Gly Gly Glu  
20 25 30

Gln Asn Gly Gln Glu Glu Lys Trp Cys Glu Lys Ala Val Lys Ser Leu

35

40

45

Val Lys Lys Leu Lys Lys Thr Gly Arg Leu Asp Glu Leu Glu Lys Ala

50

55

60

Ile Thr Thr Gln Asn Cys Asn Thr Lys Cys Val Thr Ile Pro Ser Thr

65

70

75

80

Cys Ser Glu Ile Trp Gly Leu Ser Thr Pro Asn Thr Ile Asp Gln Trp

85

90

95

Asp Thr Thr Gly Leu Tyr Ser Phe Ser Glu Gln Thr Arg Ser Leu Asp

100

105

110

Gly Arg Leu Gln Val Ser His Arg Lys Gly Leu Pro His Val Ile Tyr

115

120

125

Cys Arg Leu Trp Arg Trp Pro Asp Leu His Ser His His Glu Leu Lys

130

135

140

Ala Ile Glu Asn Cys Glu Tyr Ala Phe Asn Leu Lys Lys Asp Glu Val

145

150

155

160

Cys Val Asn Pro Tyr His Tyr Gln Arg Val Glu Thr Pro Val Leu Pro

165

170

175

Pro Val Leu Val Pro Arg His Thr Glu Ile Leu Thr Glu Leu Pro Pro

180

185

190

Leu Asp Asp Tyr Thr His Ser Ile Pro Glu Asn Thr Asn Phe Pro Ala

195

200

205

Gly Ile Glu Pro Gln Ser Asn Tyr Ile Pro Glu Thr Pro Pro Pro Gly

210

215

220

Tyr Ile Ser Glu Asp Gly Glu Thr Ser Asp Gln Gln Leu Asn Gln Ser

225

230

235

240

Ser Thr Pro Cys Trp Ile Glu Leu His Leu Asn Gly Pro Leu Gln Trp

435

440

445

Leu Asp Lys Val Leu Thr Gln Met Gly Ser Pro Ser Val Arg Cys Ser

450

455

460

Ser Met Ser

465

&lt;210&gt; 5

&lt;211&gt; 425

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 5

Met Ser Ser Ile Leu Pro Phe Thr Pro Pro Ile Val Lys Arg Leu Leu

1

5

10

15

Gly Trp Lys Lys Gly Glu Gln Asn Gly Gln Glu Glu Lys Trp Cys Glu

20

25

30

Lys Ala Val Lys Ser Leu Val Lys Lys Leu Lys Lys Thr Gly Gln Leu

35

40

45

Asp Glu Leu Glu Lys Ala Ile Thr Thr Gln Asn Val Asn Thr Lys Cys

50

55

60

Ile Thr Ile Pro Arg Ser Leu Asp Gly Arg Leu Gln Val Ser His Arg

65

70

75

80

Lys Gly Leu Pro His Val Ile Tyr Cys Arg Leu Trp Arg Trp Pro Asp

85

90

95

Leu His Ser His His Glu Leu Arg Ala Met Glu Leu Cys Glu Phe Ala

100

105

110

Phe Asn Met Lys Lys Asp Glu Val Cys Val Asn Pro Tyr His Tyr Gln

115	120	125
Arg Val Glu Thr Pro Val Leu Pro Pro Val Leu Val Pro Arg His Thr		
130	135	140
Glu Ile Pro Ala Glu Phe Pro Pro Leu Asp Asp Tyr Ser His Ser Ile		
145	150	155
		160
Pro Glu Asn Thr Asn Phe Pro Ala Gly Ile Glu Pro Gln Ser Asn Ile		
165	170	175
Pro Glu Thr Pro Pro Pro Gly Tyr Leu Ser Glu Asp Gly Glu Thr Ser		
180	185	190
Asp His Gln Met Asn His Ser Met Asp Ala Gly Ser Pro Asn Leu Ser		
195	200	205
Pro Asn Pro Met Ser Pro Ala His Asn Asn Leu Asp Leu Gln Pro Val		
210	215	220
Thr Tyr Cys Glu Pro Ala Phe Trp Cys Ser Ile Ser Tyr Tyr Glu Leu		
225	230	235
		240
Asn Gln Arg Val Gly Glu Thr Phe His Ala Ser Gln Pro Ser Met Thr		
245	250	255
Val Asp Gly Phe Thr Asp Pro Ser Asn Ser Glu Arg Phe Cys Leu Gly		
260	265	270
Leu Leu Ser Asn Val Asn Arg Asn Ala Ala Val Glu Leu Thr Arg Arg		
275	280	285
His Ile Gly Arg Gly Val Arg Leu Tyr Tyr Ile Gly Gly Glu Val Phe		
290	295	300
Ala Glu Cys Leu Ser Asp Ser Ala Ile Phe Val Gln Ser Pro Asn Cys		
305	310	315
		320

Asn Gln Arg Tyr Gly Trp His Pro Ala Thr Val Cys Lys Ile Pro Pro  
 325 330 335

Gly Cys Asn Leu Lys Ile Phe Asn Asn Gln Glu Phe Ala Ala Leu Leu  
 340 345 350

Ala Gln Ser Val Asn Gln Gly Phe Glu Ala Val Tyr Gln Leu Thr Arg  
 355 360 365

Met Cys Thr Ile Arg Met Ser Phe Val Lys Gly Trp Gly Ala Glu Tyr  
 370 375 380

Arg Arg Gln Thr Val Thr Ser Thr Pro Cys Trp Ile Glu Leu His Leu  
 385 390 395 400

Asn Gly Pro Leu Gln Trp Leu Asp Lys Val Leu Thr Gln Met Gly Ser  
 405 410 415

Pro Ser Ile Arg Cys Ser Ser Val Ser  
 420 425

<210> 6

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 6

tcgagagcca gacaaaaagc cagacattta gccagacac

39

<210> 7

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 7

tcgagtgctt ggctaaatgt ctggcttttt gtctggctc

39

<210> 8

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 8

tcgagagaca gacaaaaaga cagacattta gacagacac

39

<210> 9

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 9

tcgagtgctt gtctaaatgt ctgtcttttt gtctgtctc

39



<210> 10

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 10

tcgagagcta cataaaaagc tacatattta gctacatac

39

<210> 11

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 11

tcgagtatgt agctaaatat gtagcttttt atgtagctc

39

<210> 12

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 12

tcgagagcca gacaaggagc cagacaagga gccagacac

39

<210> 13

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 13

ctcgagtgtc tggctccttg tctggctcct tgtctggctc

40

<210> 14

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 14

tcgagagcta cataaaaagc tacatattta gctacatac

39

<210> 15

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 15

tcgagtatgt agctaaatat gtagcttttt atgtagctc

39

<210> 16

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 16

tcgaggctgc cctaaaatgt gtattccatg gaaatgtctg cccttctctc

50

<210> 17

<211> 50

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 17

tcgagagaga agggcagaca tttccatgga atacacattt tagggcagcc

50

<210> 18

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 18

ccgggatgac tcagc

15

<210> 19

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 19

ccgggctgag tcatc

15

<210> 20

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 20

ccggtttggg ttgaagccaa tatg

24

<210> 21

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 21

ccggcatatt ggcttcaatc caaa

24

<210> 22

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 22

tcgaggacag ggggcggagc ctc

23

<210> 23

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 23

tcgagaggct ccgccccctg tcc

23